

## **STIC Biotechnology Systems Branch**

### **RAW SEQUENCE LISTING** **ERROR REPORT**

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:**

Application Serial Number:

101517 079C

Source:

STIC

Date Processed by STIC:

11/28/06

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

**FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:**

**<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>**

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)**
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450**
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314**

Revised 01/10/06

## Raw Sequence Listing Error Summary

### ERROR DETECTED

### SUGGESTED CORRECTION

### SERIAL NUMBER:

10/517/079C

**ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE**

1  Wrapped Nucleics  
 Wrapped Aminos      The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2  Invalid Line Length      The rules require that a line **not exceed** 72 characters in length. This includes white spaces.

3  Misaligned Amino Numbering      The numbering under each 5<sup>th</sup> amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4  Non-ASCII      The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**

5  Variable Length      Sequence(s)       contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6  PatentIn 2.0 "bug"      A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s)      . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**

7  Skipped Sequences (OLD RULES)      Sequence(s)       missing. If intentional, please insert the following lines for **each** skipped sequence:  
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)  
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
 This sequence is intentionally skipped  
 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.

8  Skipped Sequences (NEW RULES)      Sequence(s)       missing. If intentional, please insert the following lines for **each** skipped sequence.  
 <210> sequence id number  
 <400> sequence id number  
 000

9  Use of n's or Xaa's (NEW RULES)      Use of n's and/or Xaa's have been detected in the Sequence Listing.  
 Per 1.823 of Sequence Rules, use of <220>-<223> is **MANDATORY** if n's or Xaa's are present.  
 In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.

10  Invalid <213> Response      Per 1.823 of Sequence Rules, the **only valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence

11  Use of <220>      Sequence(s)       missing the <220> "Feature" and associated numeric identifiers and responses.  
 Use of <220> to <223> is **MANDATORY** if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.  
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

12  PatentIn 2.0 "bug"      Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13  Misuse of n/Xaa      "n" can **only** represent a single nucleotide; "Xaa" can **only** represent a single amino acid



IFWO

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/517,079C

DATE: 11/28/2006  
TIME: 11:42:31

Input Set : N:\efs\11\_28\_06  
\10517079c\_efs\Substitute\_Sequence\_Listing\_20747\_230.txt  
Output Set: N:\CRF4\11282006\J517079C.raw

4 <110> APPLICANT: CePeP AB  
6 <120> TITLE OF INVENTION: Cell Penetrating Peptides  
9 <130> FILE REFERENCE: 20747/230  
11 <140> CURRENT APPLICATION NUMBER: US 10/517,079C  
C--> 12 <141> CURRENT FILING DATE: 2004-12-06  
14 <150> PRIOR APPLICATION NUMBER: PCT/IB03/03163  
15 <151> PRIOR FILING DATE: 2003-06-18  
17 <150> PRIOR APPLICATION NUMBER: US 60/391,788  
18 <151> PRIOR FILING DATE: 2002-06-25  
20 <150> PRIOR APPLICATION NUMBER: SE 0201863-8  
21 <151> PRIOR FILING DATE: 2002-06-18  
23 <160> NUMBER OF SEQ ID NOS: 31933  
25 <170> SOFTWARE: FastSEQ for Windows Version 4.0

## ERRORED SEQUENCES

38551 <210> SEQ ID NO: 2739  
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38553 <212> TYPE: PRT  
38554 <213> ORGANISM: Homo Sapiens  
38556 <220> FEATURE:  
38557 <222> LOCATION: ( 259 )...( 275 )  
38558 <223> OTHER INFORMATION: 5H6\_HUMAN

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38560 His Ser Arg Lys Ala Leu Lys Ala Ser Leu Thr Leu Gly Ile Leu Leu  
E--> 38561 1 5 10 15

E--&gt; 38565 &lt;210&gt; SEQ ID NO: 2740

38566 &lt;211&gt; LENGTH: 16

38567 &lt;212&gt; TYPE: PRT

38568 &lt;213&gt; ORGANISM: Homo Sapiens

38570 &lt;220&gt; FEATURE:

38571 &lt;222&gt; LOCATION: ( 260 )...( 276 )

38572 &lt;223&gt; OTHER INFORMATION: 5H6\_HUMAN

W--&gt; 38573 &lt;400&gt; SEQUENCE: 2740

38574 Ser Arg Lys Ala Leu Lys Ala Ser Leu Thr Leu Gly Ile Leu Leu Gly

E--&gt; 38575 1 5 10 15

38579 &lt;210&gt; SEQ ID NO: 2741

38580 &lt;211&gt; LENGTH: 16

38581 &lt;212&gt; TYPE: PRT

38582 &lt;213&gt; ORGANISM: Homo Sapiens

38584 &lt;220&gt; FEATURE:

38585 &lt;222&gt; LOCATION: ( 261 )...( 277 )

Does Not Comply  
Corrected Diskette Needed

(pg. 1-6)

See item

# 3 on

error  
summary  
sheet.

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/517,079C

DATE: 11/28/2006  
TIME: 11:42:33

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Output Set: N:\CRF4\11282006\J517079C.raw

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38596 <213> ORGANISM: Homo Sapiens  
38598 <220> FEATURE:  
38599 <222> LOCATION: ( 108 )...( 124 )  
38600 <223> OTHER INFORMATION: 5H7\_HUMAN  
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38628 <223> OTHER INFORMATION: 5H7\_HUMAN  
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38641 <222> LOCATION: ( 305 )...( 321 )  
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38652 <213> ORGANISM: Homo Sapiens  
38654 <220> FEATURE:

Same errors



See item #3

on error

Summary

Sheet

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PATENT APPLICATION: US/10/517,079C

DATE: 11/28/2006  
TIME: 11:42:33

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\10517079c\_efs\Substitute\_Sequence\_Listing\_20747\_230.txt  
Output Set: N:\CRF4\11282006\J517079C.raw

38655 <222> LOCATION: ( 248 )...( 264 )  
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38669 <222> LOCATION: ( 261 )...( 277 )  
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38683 <222> LOCATION: ( 262 )...( 278 )  
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38697 <222> LOCATION: ( 268 )...( 284 )  
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38712 <223> OTHER INFORMATION: A1AB\_HUMAN  
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38719 <210> SEQ ID NO: 2751  
38720 <211> LENGTH: 16  
38721 <212> TYPE: PRT  
38722 <213> ORGANISM: Homo Sapiens

SAME  
errors

See item

#3 or

error

Summary

Sheet

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/517,079C

DATE: 11/28/2006  
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Input Set : N:\efs\11\_28\_06  
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Output Set: N:\CRF4\11282006\J517079C.raw

38724 <220> FEATURE:  
38725 <222> LOCATION: ( 152 )...( 168 )  
38726 <223> OTHER INFORMATION: A1AB\_HUMAN  
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38733 <210> SEQ ID NC: 2752  
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38735 <212> TYPE: PRT  
38736 <213> ORGANISM: Homo Sapiens  
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38739 <222> LOCATION: ( 154 )...( 170 )  
38740 <223> OTHER INFORMATION: A1AB\_HUMAN  
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38753 <222> LOCATION: ( 155 )...( 171 )  
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38789 <210> SEQ ID NO: 2756  
38790 <211> LENGTH: 16  
38791 <212> TYPE: PRT

SAME  
errors

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/517,079C

DATE: 11/28/2006  
TIME: 11:42:33

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E--> 38855 1 5 10 15  
38859 <210> SEQ ID NO: 2761  
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Same errors

See item

#3 on

error

summary

sheet

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/517,079C

DATE: 11/28/2006  
TIME: 11:42:33

Input Set : N:\efs\11\_28\_06  
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Output Set: N:\CRF4\11282006\J517079C.raw

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38903 <212> TYPE: PRT  
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38915 <210> SEQ ID NO: 2765  
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38922 <223> OTHER INFORMATION: A1AD\_HUMAN  
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E--> 38925 1 5 10 15  
38929 <210> SEQ ID NO: 2766

SAME  
errors

The type of errors shown exist throughout  
the Sequence Listing. Please check subsequent  
sequences for similar errors.